

WO 99/61614

## SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.  
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AZIMZAI, Yalda

<120> HUMAN SOCS PROTEINS

<130> PF-0525 PCT

<140> To Be Ass

<140> 11

<150> 60/087,104; 09/216,006

<150> 30,000,000  
<151> 1998-05-28; 1998-12-17

<160> 18

## <170> PERL Program

<210> 1

<211> 288

<212> PRT

<213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte clone 1758450

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Tyr Ser Leu Ser Glu Arg Leu Ile Arg Thr Ile Ala Ala Ile Arg
20 30
      25
Ser Phe Pro His Asp Asn Val Glu Asp Leu Ile Arg Gly Gly Ala
35 45
      40
Asp Val Asn Cys Thr His Gly Thr Leu Lys Pro Leu His Cys Ala
50 60
      55
Cys Met Val Ser Asp Ala Asp Cys Val Glu Leu Leu Leu Glu Lys
65 75
      70
Gly Ala Glu Val Asn Ala Leu Asp Gly Tyr Asn Arg Thr Ala Leu
80 90
      85
His Tyr Ala Ala Glu Lys Asp Glu Ala Cys Val Glu Val Leu Leu
95 105
      100
Glu Tyr Gly Ala Asn Pro Asn Ala Leu Asp Gly Asn Arg Asp Thr
110 120
      115
Pro Leu His Trp Ala Ala Phe Lys Asn Asn Ala Glu Cys Val Arg
125 135
      130
Ala Leu Leu Glu Ser Gly Ala Ser Val Asn Ala Leu Asp Tyr Asn
140 150
      145

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<210> 2  
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<213> *Homo sapiens*

<220>  
<221> misc\_feature  
<223> Incyte clone 1834242

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 1 5 10 15  
 Arg Lys Gly Lys Arg Ser Ser Trp Gly Gly Thr Ala Ala Val Ala  
 20 25 30  
 Glu Leu Lys Pro Gly Arg Pro His Gln Phe Asp Trp Lys Ser Ser  
 35 40 45  
 Cys Glu Thr Trp Ser Val Ala Phe Ser Pro Asp Gly Ser Trp Phe  
 50 55 60  
 Ala Trp Ser Gln Gly His Cys Ile Val Lys Leu Ile Pro Trp Pro  
 65 70 75  
 Leu Glu Glu Gln Phe Ile Pro Lys Gly Phe Glu Ala Lys Ser Arg  
 80 85 90  
 Ser Ser Lys Asn Glu Thr Lys Gly Arg Gly Ser Pro Lys Glu Lys  
 95 100 105  
 Thr Leu Asp Cys Gly Gln Ile Val Trp Gly Leu Ala Phe Ser Pro  
 110 115 120  
 Trp Pro Ser Pro Pro Ser Arg Lys Leu Trp Ala Arg His His Pro  
 125 130 135  
 Gln Val Pro Asp Val Ser Cys Leu Val Leu Ala Thr Gly Leu Asn  
 140 145 150  
 Asp Gly Gln Ile Lys Ile Trp Glu Val Gln Thr Gly Leu Leu Leu  
 155 160 165  
 Leu Asn Leu Ser Gly His Gln Asp Val Val Arg Asp Leu Ser Phe  
 170 175 180

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Thr Pro Ser Gly Ser Leu Ile Leu Val Ser Ala Ser Arg Asp Lys  
 185 190 195  
 Thr Leu Arg Ile Trp Asp Leu Asn Lys His Gly Lys Gln Ile Gln  
 200 205 210  
 Val Leu Ser Gly His Leu Gln Trp Val Tyr Cys Cys Ser Ile Ser  
 215 220 225  
 Pro Asp Cys Ser Met Leu Cys Ser Ala Ala Gly Glu Lys Ser Val  
 230 235 240  
 Phe Leu Trp Ser Met Arg Ser Tyr Thr Leu Ile Arg Lys Leu Glu  
 245 250 255  
 Gly His Gln Ser Ser Val Val Ser Cys Asp Phe Ser Pro Asp Ser  
 260 265 270  
 Ala Leu Leu Val Thr Ala Ser Tyr Asp Thr Asn Val Ile Met Trp  
 275 280 285  
 Asp Pro Tyr Thr Gly Glu Arg Leu Arg Ser Leu His His Thr Gln  
 290 295 300  
 Val Asp Pro Ala Met Asp Asp Ser Asp Val His Ile Ser Ser Leu  
 305 310 315  
 Arg Ser Val Cys Phe Ser Pro Glu Gly Leu Tyr Leu Ala Thr Val  
 320 325 330  
 Ala Asp Asp Arg Leu Leu Arg Ile Trp Ala Leu Glu Leu Lys Thr  
 335 340 345  
 Pro Ile Ala Phe Ala Pro Met Thr Asn Gly Leu Cys Cys Thr Phe  
 350 355 360  
 Phe Pro His Gly Gly Val Ile Ala Thr Gly Thr Arg Asp Gly His  
 365 370 375  
 Val Gln Phe Trp Thr Ala Pro Arg Val Leu Ser Ser Leu Lys His  
 380 385 390  
 Leu Cys Arg Lys Ala Leu Arg Ser Phe Leu Thr Thr Tyr Gln Val  
 395 400 405  
 Leu Ala Leu Pro Ile Pro Lys Lys Met Lys Glu Phe Leu Thr Tyr  
 410 415 420  
 Arg Thr Phe

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 <213> Homo sapiens

<220>  
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 <223> Incyte clone 1849725

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 Pro Ala Leu Leu Glu Ser Pro Arg Pro Glu Gly Gly Glu Pro  
 20 25 30  
 Pro Arg Pro Ser Pro Glu Glu Thr Gln Gln Cys Lys Phe Asp Gly  
 35 40 45  
 Gln Glu Thr Lys Gly Ser Lys Phe Ile Thr Ser Ser Ala Ser Asp  
 50 55 60  
 Phe Ser Asp Pro Val Tyr Lys Glu Ile Ala Ile Thr Asn Gly Cys  
 65 70 75  
 Ile Asn Arg Met Ser Lys Glu Glu Leu Arg Ala Lys Leu Ser Glu

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80	85	90
Phe Lys Leu Glu Thr Arg Gly Val Lys Asp Val Leu Lys Lys		Arg
95	100	105
Leu Lys Asn Tyr Tyr Lys Lys Gln Lys Leu Met Leu Lys Glu	Ser	
110	115	120
Asn Phe Ala Asp Ser Tyr Tyr Asp Tyr Ile Cys Ile Ile Asp	Phe	
125	130	135
Glu Ala Thr Cys Glu Glu Gly Asn Pro Pro Glu Phe Val His	Glu	
140	145	150
Ile Ile Glu Phe Pro Val Val Leu Leu Asn Thr His Thr Leu	Glu	
155	160	165
Ile Glu Asp Thr Phe Gln Gln Tyr Val Arg Pro Glu Ile Asn	Thr	
170	175	180
Gln Leu Ser Asp Phe Cys Ile Ser Leu Thr Gly Ile Thr Gln	Asp	
185	190	195
Gln Val Asp Arg Ala Asp Thr Phe Pro Gln Val Leu Lys Lys	Val	
200	205	210
Ile Asp Trp Met Lys Leu Lys Glu Leu Gly Thr Lys Tyr Lys	Tyr	
215	220	225
Ser Leu Leu Thr Asp Gly Ser Trp Asp Met Ser Lys Phe Leu	Asn	
230	235	240
Ile Gln Cys Gln Leu Ser Arg Leu Lys Tyr Pro Pro Phe Ala	Lys	
245	250	255
Lys Trp Ile Asn Ile Arg Lys Ser Tyr Gly Asn Phe Tyr Lys	Val	
260	265	270
Pro Arg Ser Gln Thr Lys Leu Thr Ile Met Leu Glu Lys Leu	Gly	
275	280	285
Met Asp Tyr Asp Gly Arg Pro His Cys Gly Leu Asp Asp Ser	Lys	
290	295	300
Asn Ile Ala Arg Ile Ala Val Arg Met Leu Gln Asp Gly Cys	Glu	
305	310	315
Leu Arg Ile Asn Glu Lys Met His Ala Gly Gln Leu Met Ser	Val	
320	325	330
Ser Ser Ser Leu Pro Ile Glu Gly Thr Pro Pro Pro Gln Met	Pro	
335	340	345
His Phe Arg Lys		

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<211> 355  
<212> PRT  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<223> Incyte clone 2547840

<400> 4  
Met Ala Arg Arg Pro Arg Asn Ser Arg Ala Trp His Phe Val Leu  
1 5 10 15  
Ser Ala Ala Arg Arg Asp Ala Asp Ala Arg Ala Val Ala Leu Ala  
20 25 30  
Gly Ser Thr Asn Trp Gly Tyr Asp Ser Asp Gly Gln His Ser Asp  
35 40 45  
Ser Asp Ser Asp Pro Glu Tyr Ser Thr Leu Pro Pro Ser Ile Pro

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50	55	60
Ser Ala Val Pro Val Thr Gly Glu Ser Phe Cys Asp Cys Ala Gly	70	75
65	85	90
Gln Ser Glu Ala Ser Phe Cys Ser Ser Leu His Ser Ala His Arg	100	105
Gly Arg Asp Cys Arg Cys Gly Glu Glu Asp Glu Tyr Phe Asp Trp	115	120
80	95	110
Val Trp Asp Asp Leu Asn Lys Ser Ser Ala Thr Leu Leu Ser Cys	125	130
Asp Asn Arg Lys Val Ser Phe His Met Glu Tyr Ser Cys Gly Thr	140	145
140	155	160
Ala Ala Ile Arg Gly Thr Lys Glu Leu Gly Glu Gly Gln His Phe	170	175
Trp Glu Ile Lys Met Thr Ser Pro Val Tyr Gly Thr Asp Met Met	185	190
185	200	205
Ser Tyr Thr Gly Leu Leu His His Lys Gly Asp Lys Thr Ser Phe	210	215
Ser Ser Arg Phe Gly Gln Gly Ser Ile Ile Gly Val His Leu Asp	220	225
225	230	235
Thr Trp His Gly Thr Leu Thr Phe Phe Lys Asn Arg Lys Cys Ile	240	245
Gly Val Ala Ala Thr Lys Leu Gln Asn Lys Arg Phe Tyr Pro Met	250	255
255	260	265
Cys Ala Ser Ala Thr Ser Leu Gln Tyr Leu Cys Cys His Arg Leu	270	275
275	280	285
Arg Gln Leu Arg Pro Asp Ser Gly Asp Thr Leu Glu Gly Leu Pro	290	295
290	305	310
Leu Pro Pro Gly Leu Lys Gln Val Leu His Asn Lys Leu Gly Trp	310	315
315	320	325
Val Leu Ser Met Ser Cys Ser Arg Arg Lys Ala Pro Val Ser Asp	330	335
335	340	340
Pro Gln Ala Ala Thr Ser Ala His Pro Ser Ser Arg Glu Pro Arg	345	355
355	350	350

<210> 5  
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<213> Homo sapiens

<220>  
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<223> Incyte clone 3071986

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1 5 10 15

Leu Arg Thr Ile Gly Glu Leu Leu Ala Pro Ala Ala Pro Phe Asp  
 20 25 30  
 Lys Lys Cys Gly Arg Glu Asn Trp Thr Val Ala Phe Ala Pro Asp  
 35 40 45  
 Gly Ser Tyr Phe Ala Trp Ser Gln Gly His Arg Thr Val Lys Leu  
 50 55 60  
 Val Pro Trp Ser Gln Cys Leu Gln Asn Phe Leu Leu His Gly Thr  
 65 70 75  
 Lys Asn Val Thr Asn Ser Ser Leu Arg Leu Pro Arg Gln Asn  
 80 85 90  
 Ser Asp Gly Gly Gln Lys Asn Lys Pro Arg Glu His Ile Ile Asp  
 95 100 105  
 Cys Gly Asp Ile Val Trp Ser Leu Ala Phe Gly Ser Ser Val Pro  
 110 115 120  
 Glu Lys Gln Ser Arg Cys Val Asn Ile Glu Trp His Arg Phe Arg  
 125 130 135  
 Phe Gly Gln Asp Gln Leu Leu Leu Ala Thr Gly Leu Asn Asn Gly  
 140 145 150  
 Arg Ile Lys Ile Trp Asp Val Tyr Thr Gly Lys Leu Leu Leu Asn  
 155 160 165  
 Leu Val Asp His Thr Glu Val Val Arg Asp Leu Thr Phe Ala Pro  
 170 175 180  
 Asp Gly Ser Leu Ile Leu Val Ser Ala Ser Arg Asp Lys Thr Leu  
 185 190 195  
 Arg Val Trp Asp Leu Lys Asp Asp Gly Asn Met Met Lys Val Leu  
 200 205 210  
 Arg Gly His Gln Asn Trp Val Tyr Ser Cys Ala Phe Ser Pro Asp  
 215 220 225  
 Ser Ser Met Leu Cys Ser Val Gly Ala Ser Lys Ala Val Phe Leu  
 230 235 240  
 Trp Asn Met Asp Lys Tyr Thr Met Ile Arg Lys Leu Glu Gly His  
 245 250 255  
 His His Asp Val Val Ala Cys Asp Phe Ser Pro Asp Gly Ala Leu  
 260 265 270  
 Leu Ala Thr Ala Ser Tyr Asp Thr Arg Val Tyr Ile Trp Asp Pro  
 275 280 285  
 His Asn Gly Asp Ile Leu Met Glu Phe Gly His Leu Phe Pro Pro  
 290 295 300  
 Pro Thr Pro Ile Phe Ala Gly Gly Ala Asn Asp Arg Trp Val Arg  
 305 310 315  
 Ser Val Ser Phe Ser His Asp Gly Leu His Val Ala Ser Leu Ala  
 320 325 330  
 Asp Asp Lys Met Val Arg Phe Trp Arg Ile Asp Glu Asp Tyr Pro  
 335 340 345  
 Val Gln Val Ala Pro Leu Ser Asn Gly Leu Cys Cys Ala Phe Ser  
 350 355 360  
 Thr Asp Gly Ser Val Leu Ala Ala Gly Thr His Asp Gly Ser Val  
 365 370 375  
 Tyr Phe Trp Ala Thr Pro Arg Gln Val Pro Ser Leu Gln His Leu  
 380 385 390  
 Cys Arg Met Ser Ile Arg Arg Val Met Pro Thr Gln Glu Val Gln  
 395 400 405  
 Glu Leu Pro Ile Pro Ser Lys Leu Leu Glu Phe Leu Ser Tyr Arg  
 410 415 420

Ile

<210> 6  
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<213> Homo sapiens

<220>  
<221> misc\_feature  
<223> Incyte clone 3484619

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Phe Trp Val Glu Arg Thr Pro Val His Glu Ala Ala Gln Arg Gly  
20 25 30  
Glu Ser Leu Gln Leu Gln Gln Leu Ile Glu Ser Gly Ala Cys Val  
35 40 45  
Asn Gln Val Thr Val Asp Ser Ile Thr Pro Leu His Ala Ala Ser  
50 55 60  
Leu Gln Gly Gln Ala Arg Cys Val Gln Leu Leu Ala Ala Gly  
65 70 75  
Ala Gln Val Asp Ala Arg Asn Ile Asp Gly Ser Thr Pro Leu Cys  
80 85 90  
Asp Ala Cys Ala Ser Gly Ser Ile Glu Cys Val Lys Leu Leu Leu  
95 100 105  
Ser Tyr Gly Ala Lys Val Asn Pro Pro Leu Tyr Thr Ala Ser Pro  
110 115 120  
Leu His Glu Ala Cys Met Ser Gly Ser Ser Glu Cys Val Arg Leu  
125 130 135  
Leu Ile Asp Val Gly Ala Asn Leu Glu Ala His Asp Cys His Phe  
140 145 150  
Gly Thr Pro Leu His Val Ala Cys Ala Arg Glu His Leu Asp Cys  
155 160 165  
Val Lys Val Leu Leu Asn Ala Gly Ala Asn Val Asn Ala Ala Lys  
170 175 180  
Leu His Glu Thr Ala Leu His His Ala Ala Lys Val Lys Asn Val  
185 190 195  
Asp Leu Ile Glu Met Leu Ile Glu Phe Gly Gly Asn Ile Tyr Ala  
200 205 210  
Arg Asp Asn Arg Gly Lys Lys Pro Ser Asp Tyr Thr Trp Ser Ser  
215 220 225  
Ser Ala Pro Ala Lys Cys Phe Glu Tyr Tyr Glu Lys Thr Pro Leu  
230 235 240  
Thr Leu Ser Gln Leu Cys Arg Val Asn Leu Arg Lys Ala Thr Gly  
245 250 255  
Val Arg Gly Leu Glu Lys Ile Ala Lys Leu Asn Ile Pro Pro Arg  
260 265 270  
Leu Ile Asp Tyr Leu Ser Tyr Asn  
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<210> 7  
<211> 281  
<212> PRT  
<213> Homo sapiens

<220>  
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<223> Incyte clone 1275743

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Lys Phe Leu Leu Val Gly Asp Ser Asp Val Gly Lys Gly Glu Ile  
20 25 30  
Leu Glu Ser Leu Gln Asp Gly Ala Ala Glu Ser Pro Tyr Ala Tyr  
35 40 45  
Ser Asn Gly Ile Asp Tyr Lys Thr Thr Ile Leu Leu Asp Gly  
50 55 60  
Arg Arg Val Lys Leu Glu Leu Trp Asp Thr Ser Gly Gln Gly Arg  
65 70 75  
Phe Cys Thr Ile Phe Arg Ser Tyr Ser Arg Gly Ala Gln Gly Ile  
80 85 90  
Leu Leu Val Tyr Asp Ile Thr Asn Arg Trp Ser Phe Asp Gly Ile  
95 100 105  
Asp Arg Trp Ile Lys Glu Ile Asp Glu His Ala Pro Gly Val Pro  
110 115 120  
Arg Ile Leu Val Gly Asn Arg Leu His Leu Ala Phe Lys Arg Gln  
125 130 135  
Val Pro Thr Glu Gln Ala Arg Ala Tyr Ala Glu Lys Asn Cys Met  
140 145 150  
Thr Phe Phe Glu Val Ser Pro Leu Cys Asn Phe Asn Val Ile Glu  
155 160 165  
Ser Phe Thr Glu Leu Ser Arg Ile Val Leu Met Arg His Gly Met  
170 175 180  
Glu Lys Ile Trp Arg Pro Asn Arg Val Phe Ser Leu Gln Asp Leu  
185 190 195  
Cys Cys Arg Ala Ile Val Ser Cys Thr Pro Val His Leu Ile Asp  
200 205 210  
Lys Leu Pro Leu Pro Val Thr Ile Lys Ser His Leu Lys Ser Phe  
215 220 225  
Ser Met Ala Asn Gly Met Asn Ala Val Met Met His Gly Arg Ser  
230 235 240  
Tyr Ser Leu Ala Ser Gly Ala Gly Gly Ser Lys Gly Asn  
245 250 255  
Ser Leu Lys Arg Ser Lys Ser Ile Arg Pro Pro Gln Ser Pro Pro  
260 265 270  
Gln Asn Cys Ser Arg Ser Asn Cys Lys Ile Ser  
275 280

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<213> Homo sapiens

<220>  
<221> misc\_feature  
<223> Incyte clone 1722533

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Gln Glu Glu Tyr Ser Leu Tyr Ser Ser Leu Ser Glu Asp Glu Leu  
 20 25 30  
 Val Gln Met Ala Ile Glu Gln Ser Leu Ala Asp Lys Thr Arg Gly  
 35 40 45  
 Pro Thr Thr Ala Glu Ala Thr Ala Ser Ala Cys Thr Asn Arg Gln  
 50 55 60  
 Pro Ala His Phe Tyr Pro Trp Thr Arg Ser Thr Ala Pro Pro Glu  
 65 70 75  
 Ser Ser Pro Ala Arg Ala Pro Met Gly Leu Phe Gln Gly Val Met  
 80 85 90  
 Gln Lys Tyr Ser Ser Ser Leu Phe Lys Thr Ser Gln Leu Ala Pro  
 95 100 105  
 Ala Asp Pro Leu Ile Lys Ala Ile Lys Asp Gly Asp Glu Glu Ala  
 110 115 120  
 Leu Lys Thr Met Ile Lys Glu Gly Lys Asn Leu Ala Glu Pro Asn  
 125 130 135  
 Lys Glu Gly Trp Leu Pro Leu His Glu Ala Ala Tyr Tyr Gly Gln  
 140 145 150  
 Val Gly Cys Leu Lys Val Leu Gln Arg Ala Tyr Pro Gly Thr Ile  
 155 160 165  
 Asp Gln Arg Thr Leu Gln Glu Glu Thr Ala Val Tyr Leu Ala Thr  
 170 175 180  
 Cys Arg Gly His Leu Asp Cys Leu Leu Ser Leu Leu Gln Ala Gly  
 185 190 195  
 Ala Glu Pro Asp Ile Ser Asn Lys Ser Arg Glu Thr Pro Leu Tyr  
 200 205 210  
 Lys Ala Cys Glu Arg Lys Asn Ala Glu Ala Val Lys Ile Leu Val  
 215 220 225  
 Gln His Asn Ala Asp Thr Asn His Arg Cys Asn Arg Gly Trp Thr  
 230 235 240  
 Ala Leu His Glu Ser Val Ser Arg Asn Asp Leu Glu Val Met Gln  
 245 250 255  
 Ile Leu Val Ser Gly Gly Ala Lys Val Glu Ser Lys Asn Ala Tyr  
 260 265 270  
 Gly Ile Thr Pro Leu Phe Val Ala Ala Gln Ser Gly Gln Leu Glu  
 275 280 285  
 Ala Leu Arg Phe Leu Ala Lys Tyr Gly Ala Asp Ile Asn Thr Gln  
 290 295 300  
 Ala Ser Asp Asn Ala Ser Ala Leu Tyr Glu Ala Cys Lys Asn Glu  
 305 310 315  
 His Glu Glu Val Val Glu Phe Leu Leu Ser Gln Gly Ala Asp Ala  
 320 325 330  
 Asn Lys Thr Asn Lys Asp Gly Leu Leu Pro Leu His Ile Ala Ser  
 335 340 345  
 Lys Lys Gly Asn Tyr Arg Ile Val Gln Met Leu Leu Pro Val Thr  
 350 355 360  
 Ser Arg Thr Arg Ile Arg Arg Ser Gly Val Ser Pro Leu His Leu  
 365 370 375  
 Ala Ala Glu Arg Asn His Asp Glu Val Leu Glu Ala Leu Leu Ser  
 380 385 390  
 Ala Arg Phe Asp Val Asn Thr Pro Leu Ala Pro Glu Arg Ala Arg  
 395 400 405  
 Leu Tyr Glu Asp Arg Arg Thr Ser Ala Leu Tyr Phe Ala Val Val  
 410 415 420  
 Asn Asn Asn Val Tyr Ala Thr Glu Leu Leu Leu Gln His Gly Ala  
 425 430 435  
 Asp Pro Asn Arg Asp Val Ile Ser Pro Leu Leu Val Ala Ile Arg

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440	445	450
His Gly Cys Leu Arg Thr Met Gln Leu Leu Leu Asp His Gly Ala		
455	460	465
Asn Ile Asp Ala Tyr Ile Ala Thr His Pro Thr Ala Phe Pro Ala		
470	475	480
Thr Ile Met Phe Ala Met Lys Cys Leu Ser Leu Leu Lys Phe Leu		
485	490	495
Met Asp Leu Gly Cys Asp Gly Glu Pro Cys Phe Ser Cys Leu Tyr		
500	505	510
Gly Asn Gly Pro His Pro Pro Ala Pro Gln Pro Ser Ser Arg Phe		
515	520	525
Asn Asp Ala Pro Ala Ala Asp Lys Glu Pro Ser Val Val Gln Phe		
530	535	540
Cys Glu Phe Val Ser Ala Pro Glu Val Ser Arg Trp Ala Gly Pro		
545	550	555
Ile Ile Asp Val Leu Leu Asp Tyr Val Gly Asn Val Gln Leu Cys		
560	565	570
Ser Arg Leu Lys Glu His Ile Asp Ser Phe Glu Asp Trp Ala Val		
575	580	585
Ile Lys Glu Lys Ala Glu Pro Pro Arg Pro Leu Ala His Leu Cys		
590	595	600
Arg Leu Arg Val Arg Lys Ala Ile Gly Lys Tyr Arg Ile Lys Leu		
605	610	615
Leu Asp Thr Leu Pro Leu Pro Gly Arg Leu Ile Arg Tyr Leu Lys		
620	625	630
Tyr Glu Asn Thr Gln		
635		

<210> 9  
<211> 518  
<212> PRT  
<213> Homo sapiens

<220>  
<221> misc\_feature  
<223> Incyte clone 1759763

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1	5	10
Leu Ala Ala Arg Glu Gly Asn Val Lys Val Leu Arg Lys Leu Leu		
20	25	30
Lys Lys Gly Arg Ser Val Asp Val Ala Asp Asn Arg Gly Trp Met		
35	40	45
Pro Ile His Glu Ala Ala Tyr His Asn Ser Val Glu Cys Leu Gln		
50	55	60
Met Leu Ile Asn Ala Asp Ser Ser Glu Asn Tyr Ile Lys Met Lys		
65	70	75
Thr Phe Glu Gly Phe Cys Ala Leu His Leu Ala Ala Ser Gln Gly		
80	85	90
His Trp Lys Ile Val Gln Ile Leu Leu Glu Ala Gly Ala Asp Pro		
95	100	105
Asn Ala Thr Thr Leu Glu Glu Thr Thr Pro Leu Phe Leu Ala Val		
110	115	120

Glu Asn Gly Gln Ile Asp Val Leu Arg Leu Leu Leu Gln His Gly  
 125 130 135  
 Ala Asn Val Asn Gly Ser His Ser Met Cys Gly Trp Asn Ser Leu  
 140 145 150  
 His Gln Ala Ser Phe Gln Glu Asn Ala Glu Ile Ile Lys Leu Leu  
 155 160 165  
 Leu Arg Lys Gly Ala Asn Lys Glu Cys Gln Asp Asp Phe Gly Ile  
 170 175 180  
 Thr Pro Leu Phe Val Ala Ala Gln Tyr Gly Lys Leu Glu Ser Leu  
 185 190 195  
 Ser Ile Leu Ile Ser Ser Gly Ala Asn Val Asn Cys Gln Ala Leu  
 200 205 210  
 Asp Lys Ala Thr Pro Leu Phe Ile Ala Ala Gln Glu Gly His Thr  
 215 220 225  
 Lys Cys Val Glu Leu Leu Leu Ser Ser Gly Ala Asp Pro Asp Leu  
 230 235 240  
 Tyr Cys Asn Glu Asp Ser Trp Gln Leu Pro Ile His Ala Ala Ala  
 245 250 255  
 Gln Met Gly His Thr Lys Ile Leu Asp Leu Leu Ile Pro Leu Thr  
 260 265 270  
 Asn Arg Ala Cys Asp Thr Gly Leu Asn Lys Val Ser Pro Val Tyr  
 275 280 285  
 Ser Ala Val Phe Gly Gly His Glu Asp Cys Leu Glu Ile Leu Leu  
 290 295 300  
 Arg Asn Gly Tyr Ser Pro Asp Ala Gln Ala Cys Leu Val Phe Gly  
 305 310 315  
 Phe Ser Ser Pro Val Cys Met Ala Phe Gln Lys Asp Cys Glu Phe  
 320 325 330  
 Phe Gly Ile Val Asn Ile Leu Leu Lys Tyr Gly Ala Gln Ile Asn  
 335 340 345  
 Glu Leu His Leu Ala Tyr Cys Leu Lys Tyr Glu Lys Phe Ser Ile  
 350 355 360  
 Phe Arg Tyr Phe Leu Arg Lys Gly Cys Ser Leu Gly Pro Trp Asn  
 365 370 375  
 His Ile Tyr Glu Phe Val Asn His Ala Ile Lys Ala Gln Ala Lys  
 380 385 390  
 Tyr Lys Glu Trp Leu Pro His Leu Leu Val Ala Gly Phe Asp Pro  
 395 400 405  
 Leu Ile Leu Leu Cys Asn Ser Trp Ile Asp Ser Val Ser Ile Asp  
 410 415 420  
 Thr Leu Ile Phe Thr Leu Glu Phe Thr Asn Trp Lys Thr Leu Ala  
 425 430 435  
 Pro Ala Val Glu Arg Met Leu Ser Ala Arg Ala Ser Asn Ala Trp  
 440 445 450  
 Ile Leu Gln Gln His Ile Ala Thr Val Pro Ser Leu Thr His Leu  
 455 460 465  
 Cys Arg Leu Glu Ile Arg Ser Ser Leu Lys Ser Glu Arg Leu Arg  
 470 475 480  
 Ser Asp Ser Tyr Ile Ser Gln Leu Pro Leu Pro Arg Ser Leu His  
 485 490 495  
 Asn Tyr Leu Leu Tyr Glu Asp Val Leu Arg Met Tyr Glu Val Pro  
 500 505 510  
 Glu Leu Ala Ala Ile Gln Asp Gly  
 515

097023-070502

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<210> 10  
<211> 1117  
<212> DNA  
<213> *Homo sapiens*

<220>  
<221> misc\_feature  
<223> Incyte clone 1758450

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 cacgccttga cagcggctt caaccccccac ctcagccccag caattcggca gtttggagca 60  
 tgtgaacacc ttgagcccttg atgagttcca gtatgtggta tattatgcag agcattcaga 120  
 gcaaataactc tctctcccgag cgcttaatcc gaacaattgc tgccatccgt tccttcccac 180  
 atgataatgt agaggacctc atcagagggg gagcagatgt gaaactgcact catggcacac 240  
 tgaagccctt gcactgtgcc tgtatggtgt cagatgtcga ctgtgtggag ttacttctgg 300  
 aaaaaggagc cgaggtgaat gcccctggatg ggtataaccg aacagccctc cactatgcag 360  
 cagagaaga tgaggcttgt gtggagggtcc tattggagta tggtgcaaac cccaaatgcctt 420  
 tggatggcaa cagagatacc ccacttcaact gggcagccctt taagaacaat gctgagtg 480  
 tgcgggctct cctagagagc gggggcctctg tcaatgcctt ggattacaac aatgatacac 540  
 cgctcagctg ggctgccatg aaggggaaatc ttgagagtgt cagcatccctt ctggattatg 600  
 ggcgcagaggt cagagtcatc aacctaatacg gccagacacc catctccgc ctgggtggctc 660  
 tgcttagtcag gggacttggaa acagagaaaag aggactcttg ctttgagctc ctccacagag 720  
 ctgttggaca ctttgaatttggaaaatg gcacccatgcc acgagaggtg gccagagacc 780  
 cgcagctatg tgaaaaaaactg actgttctgt gctcagctcc aggaactcta aaaacactcg 840  
 ctcgctatgc cgtgcgccgt agcctgggac tccagtatct ccccgatgca gtgaaggggcc 900  
 ttccactgccc agtttctttt aaggaataacc ttttactttt agaatacgcc gagaagatgt 960  
 ttgcaccatc gtgcaggccag ctctgggtga ggttgccttgcagttactcc ttgtcaca 1020  
 aaacagaaaa acagttgttt cctgtatgtgt gggttataga tttcgaagca acatgtcaca 1080  
 acaataaacct gcatagcaac tcccccttcc aaacaaaa 1117

<210> 11  
<211> 2589  
<212> DNA  
<213> *Homo sapiens*

<220>  
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<223> Incyte clone 1834242

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